Nucleotide sequence of SSL2 genomic gene (wild type)	
ATGAGTAGTTTGGTGGAGAGGCTTCGCATACGATCTGATAGGAAACCAGTTTATAACCTAGATGATTCTGATGATG	9 /
A C G A C T T C G T T C C T A A A A A A A G A C C T T T G A G C A A G T C G A G G C T A T T G T C A G A A C T G G G L L L G L L C L C	152
ctctcgagcttattgttcagcttttactgttttatgtgttctattttaatgtgtattttagtctttttttgtgttgttactctgaa	228
tttgtagaaagaaangcangncagccnngngggaaagnacnacnngnagcngcaanacangcacniangcg	304
TTCCATGCTAAATGCTTAGTTCCACCTCTTAAAGATGCTTCCGTGGAAAATTGGAGATGCCCTGAATGTgtaagat	3 8 0
tttagttacggtccacaattatgttttgggatgctacaggttccatttttttt	456
t t t g c a g G T T A G T C C T C T T A A C G A G A T A G A T A A G A T T G G A T T G T G	532
GGTTCCTCCGATGCGGAACCGAAGCCAATTTTTGTGAAACAGTATCTCGTGAAGTGGAAGGGATTATCATACCTTC	809
ACTGCTCTTGgtagttactgcgtgtcttttttgctgtctggcacgctaattatcatgtttctttc	684
tataatatgtgatttatttccttttactaatcatag6GTGCCTGAGAAGGAGTTCCAGAAGGCTTATAAGTCAAAT	092
CATCGTTTAAAAACCAGAGTGAACAATTTTCACCGTCAAATGGAGTCATTCAATAACAGGAGAGATGATTTGTTG	836
CCATACGTCCTGAGTGGACCACTGTTGATCGGATTCTTGCCTGCAGgtctagagaatggaattaattcctttattt	912
atctatctgccaactttttttttaatatccttgttttcagcataatccatctattctattctctaataacacgtatcttgata	8 8 6
gagtgetgettaacetaaatttaetgttateaegattttgggtetetgaaaeatgataaatgaeetgeetg	1064
tttttcttctttttaagttaccattttcttagttgtttcgtaaatcaggaattgtgacagttgcattggtttcttt	1140
tatgatatagaGAGGAAGATGGGGAGCTGGAATATCTTGTCAAATATAAAGAGCTATCCTATGATGAATGTTATTG	1216
G G A G T C A G A A T C A G A C C T T C C A G A A T G A A A T T C A A A G G T T C A A G G A T G T A A A T T C T A G A A C T C G C A G A	1292

<u>ئ</u>

A GTA A A GATGTT G A C C A T A A A A A A A A A A C C A G A G A C T T C A A G A T T G A T C C T G A A T T C C T C A A A G G t a	1368
tttggatcaccttaaaatcatatactataaatgtttcttatatttggtacttatagatgttatgatttattt	1444
	1520
TGTAATCC	1596
catccattgtaaggggcctttgttttctattcctgtaatgttgtgagatttttcctgttacagGCAAGACAATTCA	1672
A A G C A T T G C C C T T T T T C A C T T T T T G A G A A A C C T C A T T C C G C A T T T G G T A A T T G C T C T C T A T C G A C T C T G	1748
CGTAACTGGGAGAGAGTTTGCCACATGGGCCCCACAGATGAACGTGgtatgtatgcagttatacacgcaatgat	1824
ctgtgccatttgtatgttttttgttgtttgtttattggttaatggaatggtcttcgtggtcatttgacgggtagGTTATGTATTT	1900
CACTGCGC	1976
A A G A A A T C T G G A C A A A T A A G C G A A A G C A A A A A A A A A G A A T C A G T T T G A T G C T C C T C A C A T C G T A T G A G A	2025
T G A T C A A C C T A G A T T C A G C A G T T C T A A A A C C A A T T A A G T G G A G T G C A T G g t a a t t c t c t a a t g a g a c t t	2128
tactttctcttagtcgtctctctttctctttacatgttgcctagtaacattgttttgggcagATTGTTGATGAA	2204
GGTCATCGACTGAAAAAAAGGATTCAAAGCTGTTCTCTTCATTGACACAGTATTCAAGTAACCACCGTATTCTTC	2280
T G A C A G G A A C A C C A C T T C A G g t t c g t c a t t t g a g t t t c t g a a g t t t c a a t a g t t g t a t c t g a g c	2356
atagtagctacgatttgcaatgagaattgttatatattatcttgcactaatgtcttacctgattagttgcaatatg	2432
tgatgat	2508
GAAGgtatcacaagaatagcaaagataaataagttcgcatacttaacagaatttatgtagctaacatgttatt	2584
attgcacaatacttgcagTTTGGAAGTTTGGAGGAGTTCCAGGAGTTCAAAGATATTAATCAAGAGAGA	2660

F | G. 2

T C T C A A G G T T G C A C A A A T G T T G G C T C C A C A T T T G C T C A G A A g t a a c c a a a a c t a t t t g t t c a t c t t t t t t a a	2736
tttatatatgtgtttcaaaagtttggttggagggaatctttcatagtaataattttatgatcttaaccatgctgtctc	2812
attactcttccag G G G T A A A A A A	2888
GTTGATCTGAGCAGTCTGC	2964
GGAGGTGCTCAAgtaagttcttttaatttttgtttacactttttggatcattaaacctcataggtggggtagaaa	3040
a a t	3116
ACGAAAAGTATGCTGCCATC	3192
CAAqtaatatctcatttcccaaaaatggttatctgtttattactacttattaaagtcgtctgctaacttttgcgtt	3268
gaacgttttcttatatgtatcaaagACAACTTTTGGAGTCTTGTGGAAAGCTGCAACTTCTAGATAAATGATGGT	3344
AAACTGAAAGAGCAA	3420
TGTACCCATAAGgtatttgaacttcttatatgtacagtctgtttcagtagattttcatttcattcttgttgtttttgtaga	3496
a c t g t	3572
agcatgtgctgaaattgagagtatatgagcattctgtgcccaactgaaagagcaaagaccacaaagtttccttataa	3648
acagtaca	3724
t q t g t a c a g a g t t c t c a t a a a c a c a c a g t t t c a g c a t a t g c t g g a t c t t a t a a g g t a	3800
gaacttgt	3876
cactc	3952
g caattacttt cttagaatttttg t cattacttact cigttgg caatataactt ctttatt ccct caaagattact	4028

ttttttggtttcttgaaatgccattatcaataccattgcttttgctgacgcatgcacttgagacaacttgtttta	4104
ctagcac	4180
ctttgtcagAAATGGCAGTACGAGCGAATTGATGGAAAGGTTGGCGGAGCTGAGCGGCAAATACGCATAGATCGGT	4256
000	4332
GGCTGATACAGTAATCATTTATGACAGgtttgaatttcagcttctcttagtgtcatctgtactctttcatagtta	4408
ttgtgtcaagctgtaagaggaactatttggcttgatagcataatattggatagcataatatttggaagtttaatgttgattttaagtga	4484
gggttgtgatgagtga	4560
D	4636
cgtcca	4712
ctgttttatc	4 7 8 8
CCAAACAAATAAGgttttaaaattttatctcttagtgctgtcaacttgcaattttgtgttctttttgtagtttccc	4864
taattttccttatattttcctttagGTGATGATTTATAGGCTCATAAACCGAGGCACCATTGAAGAAAGGATGATG	4940
CAATTGACTAAAAAGAAAATGGTTCTAGAGCATCTTGTTGTTGGGAAACTCAAAACACAAAAAAATTAATCAGgtaa	5016
actittattgcttgaagcctttttacttgattacaaatttctcaacggattggagctggaaggtagaaattccaag	2605
a a ga a cacctt c ggttata a cttata a gtgtga a atta a a a gata a a a a	5168
ttgtttgtca	5244
ATC	5320
GCTATAGACAAgtaatagactccttactctttcctcttgttttgttttg	5396

F. G. 4

qattgctcctttcttatgaaagcttttgcagtcaattgcatgggcgtatttcatttttgtctctctatctgtctctgtct	5472
q c a g A T T G C T T G A T C G T G A T C T C G T G G A G	5 5 4 8
ر و	5624
ggagtct	5700
ctgctgtg	5776
ىد ب	5852
ttagtgatgatga	5928
tttgaaccctgatc	6004
tgcttg	0809
gttctgaagcaaatag	6156
atcttcc	6232
CGCTGCTGAAAGCAAA	6308
TTGAGCTGCACCAGGCTGAGGAGCTTAATGCTCTTGGAAAAAGGAAGAGAAGAGAAGGAAG	6384
t gat c c c c c t t a t c c a a t t g t g g c a t c a t a t t g a t a a c t g g a t t t t t c a c c a t t t a t g t c t t c t g a t t c t g t c c	6460
tgtttcatatatttattcatgttgtctaacttttccttttgaattccttaggtagctaaattcagaaagtaataat	6536
ttagttgactgtatccttctaaattgagaaagtataatttagttgactgtatccagtataaaactaaacgcccttg	6612
Ü	8 8 9 9
ttaactatagtettetgttteetetgeaagaaataegttttgttteaetetetaaettgatatagatatageteaattaet	6764

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atatacattggtttggtctgccatcatcgtttcatgtctttagtguttcaatgaggctgttctaattag	6840
GGTATCCAT	6916
G A A G C T G A G T C A A C A G A T G G T G A A G C A G G A C A A G G A G T T C A G A C G G G T C G A C G T A C A G A A G A A	6992
G C G g t a t t a c c a c g t t t c g g a t t t a a t t t a a t t t g t a a t g g a g c t g a a t g a t a t a g	2068
tagatgagt	7144
CTGGGTTT	7220
t a g a	7296
tcaatattt	7372
TTTGTTCCTCGCTTAAAGCAGAAGACTT	7448
c t g	7524
CAGATATGGAATACTC	A 7600
caata	t 7676
attaatataatggtegttgt	C 7752
GCTCTTCTGATACTAGTTCAGGAGA	a 7828
aatttttttagtatatgttacttctag	t 7904
tcaaacctcactaagatt	T 7980
AACCT	A 8056
G G A A G G A A C A T G A C A A G A T A A T G A T A C G T G C T G T T T T A A B t a t g a a c c c t g c a c c a c t g t t c t t a c	c 8132

a a a ta a t t t t a t t t t c t c a t t c t c	8 2 0 8
GCATGGGTACGGACGGTGGCAAGCTATTGTTGATGACAAAGAGTTGGGGATCCAAGAGCTTATCTGCAAAGAT	2 8
GAATTICCCTCACATAAGTTTGTCTGCTGCTGAACAAGCTGGTTTGCAGGGGCAGAATGGTAGTGGGGGGGCTCTAAT	8360
C C G G G A G C A C A G A C T A A C C A G A A T C C T G G A A G C G T T A T T A C T G G A A C A A T A A T G C T T C T G C T G A T G G G C T C A A G	8436
T A A A C T C G A T G T T C T A T T T C G G G A C A T G C A G A C G A C T T G T T G A G T T G T G A A A A A G C G A G T T C T G C T T T T G G A	8512
GAAGGCGATGAATTATGAATACGCAGAGGAATATTATgtatgttgtaccatctgcagtgttggtacttactcacat	8 5 8 8
gttttgcgctgaattgtttaactttgattgaatctctggttgcagGGACTTGGTGGCCTCATCATCTATCCTACTG	8664
A A G A A C C A G A A G C T G A A C A A A G A T C G C T G A C A C A G T G G G C T T T A T T G A G G T T G A T G A T G A A T G C T T G A	8740
T G G A C T T C C T A A G A C T G A T C C T A T C A g t a a g t t c c a t c a c a a g t t t t a a c g a g t t g t t c t a a t g t	8816
g a g c t c t c t g a a t c t c g c t g c a g C T T C A G A A A A T T A T G G G G C T G C T G T T G A C A A C A A C C A A G C G G G G T C G A A	8892
ATAGCTCAACATTATAACCAGgtaagctatgctttttcctttggtggtaggctaatgtctagaactagtatatca	8968
cactaatatctctccggttattcagATGTGCAAACTTCTTGATGAGAACGCTCGGGAATCAGTCCAAGCATATGTA	9044
A A C A A C C A A C C A C C G A G T A C C A A G G T G A A T G A G C T T C C G T G C A C T C A A T C A A T G G T A A C A T T A A C A C A A	9120
T C C T T T C G A T T A C A T C T G A T C C A A G T C A C A T G A A G A C G A C A C C A G A C C T A A A C A T G T T G A G A T G A A	9196
GGACACGGCCGAAGAAACAAAACCGTTAAGAGGTGGCGTCGTCGATCTGAATGTGGGGGGGG	9272
GCTGAAGCTAGTGGAAGTGTTGATGTAAAATGGAAGAAGCCAAAGAAGAAGAAGAAGAGGAGGAGAGAAGA	9348
ATTGA	9353

Г С.

Nucleotide sequence of mutant IAA14 gene	
A T G A A C C T T A A G G A G A G C G T T T G T C T T G G C C T C C C G G A G G C A C T G A A A C C G G T C G G C C A A G T C G G	9 2
GTGTTGGGAACAAGAGAGGCTTCTCCGAGACCGTTGATCTCAAACTTAATCTTCAATCTAACAAACA	152
GGATCTCAACACTAATGGAGCTCCCAAGGAGAAGACCTTCCTT	228
ctatttacacaattccttaagaagaaccttccttaaagggaaggcctttttttt	304
actaatagttgatataaaagttcttaaaatacatatatat	380
a a t g t g t g t g t g t t a t a t t c t a t g c a g A G C A C A A G T G G T T G G C C A T C G G T G A G G A A C T A C C G G A A A A	456
A T G T T A T G G C T A A T C A G A A G C G G C G A A G C A G A G G C A A T G A G T G G T G G A G G A A C C G T C G C C T T G T G A A	532
G G T T T C C A T G G A T G G A G C T C C T T A T C T T C G G A A G G T T G A C C T C A G A T C C T C T C T C T C T G A T	809
GCCTTGGCCAAAATGTTCAGCTCCTTTACCATGGgtatgcattttcagacatataagtcgaattatcattatt	684
tttgtgtttacttacaattttttttttaacgatacagtttttttcattttta	092
gggatttgattaattaagGGAGTTATGGAGCACAAGGGATGATAGATTTCATGAACGAGAGTAAAGTGATGGATC	8 3 6
T G T T G A A C A G T T C T G A G T A T G T T C C A A G C T A C G A G G A C A A G A T G G T G G A T G T G G T G A T G T C C C C T G	912
GCCgtgagtttcctcattcttcttgctttcattattatgaccaaaattattctctctaaacaaaaaaacaatattct	9 8 8
ctaaagcattattattgatattacttatcaaaaaatacacaaaatgataatcaatgtatctatgtgttataaacacg 1	1064
cacagecatettttggttggeatgggacagaaeteagagacagagaeagagatgtttatatatataaataetaaeteatea	1140
atatgttacctcatttgtagctggcacatattctttcactttcaatagatttctaaattgtcaccaacccaaat 1	1216
cccgatttcagGATGTTTGTCGAGTCATGCAAACGTTTGCGCATAATGAAAGGATCCGAAGCAATTGGACTTGgta 1	1292
agttttcttttctgttcgtttctataagtggctcttttctgttttccaataatgctcgtgtttttttt	1368
CAAGAGCAATGGAGAAGTTCAAGAACAGATCATGA	1403